p#18



1042

ENTERED

600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/581,651A

DATE: 04/12/2002 TIME: 14:21:03

Input Set : A:\350013-72.txt

```
4 <110> APPLICANT: Schor, Seth Laurence
         Schor, Ana Maria
 7 <120> TITLE OF INVENTION: POLYPEPTIDES, POLYNUCLEOTIDES AND USES
10 <130> FILE REFERENCE: 350013-72
12 <140> CURRENT APPLICATION NUMBER: 09/581,651A
13 <141> CURRENT FILING DATE: 2000-10-10
                                                            TECH CENTER 1600/2900
15 <160> NUMBER OF SEQ ID NOS: 15
17 <170> SOFTWARE: FastSEQ for Windows Version 3.0
19 <210> SEQ ID NO: 1
20 <211> LENGTH: 660
21 <212> TYPE: PRT
22 <213> ORGANISM: Human
24 <400> SEQUENCE: 1
   Asn Leu Val Ala Thr Cys Leu Pro Val Arg Ala Ser Leu Pro His Arg
26
                     5
    Leu Asn Met Leu Arg Gly Pro Gly Pro Gly Leu Leu Leu Ala Val
27
28
                                    25
    Gln Cys Leu Gly Thr Ala Val Pro Ser Thr Gly Ala Ser Lys Ser Lys
30
    Arg Gln Ala Gln Gln Met Val Gln Pro Gln Ser Pro Val Ala Val Ser
31
33
    Gln Ser Lys Pro Gly Cys Tyr Asp Asn Gly Lys His Tyr Gln Ile Asn
34
                                             75
                        70
    Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys
35
    Tyr Gly Gly Ser Arg Gly Phe Asn Cys Glu Ser Lys Pro Glu Ala Glu
37
38
                                    105
    Glu Thr Cys Phe Asp Lys Tyr Thr Gly Asn Thr Tyr Arg Val Gly Asp
39
40
            115
                                120
41
    Thr Tyr Glu Arg Pro Lys Asp Ser Met Ile Trp Asp Cys Thr Cys Ile
                            135
                                                 140
    Gly Ala Gly Arg Gly Arg Ile Ser Cys Thr Ile Ala Asn Arg Cys His
                                             155
                        150
    Glu Gly Gly Gln Ser Tyr Lys Ile Gly Asp Thr Trp Arg Arg Pro His
45
46
                                        170
    Glu Thr Gly Gly Tyr Met Leu Glu Cys Val Cys Leu Gly Asn Gly Lys
47
                                                         190
48
                                    185
                180
    Gly Glu Trp Thr Cys Lys Pro Ile Ala Glu Lys Cys Phe Asp His Ala
49
                                200
                                                     205
    Ala Gly Thr Ser Tyr Val Val Gly Glu Thr Trp Glu Lys Pro Tyr Gln
52
                            215
    Gly Trp Met Met Val Asp Cys Thr Cys Leu Gly Glu Gly Ser Gly Arg
```

RAW SEQUENCE LISTING

DATE: 04/12/2002 TIME: 14:21:03

PATENT APPLICATION: US/09/581,651A

Input Set : A:\350013-72.txt

E 4	225					230					235					240
54		ωh ν	Cvc	Thr	Sor		Asn	Δrα	Cvs	Δsn		Gln	Asp	Thr	Ara	
55 56	TTE	1111	Cys	1111	245	пту	LOH	ni 9	CyD	250		01			255	
	Cor	Фттъс	λνα	τla		λen	Thr	Ͳrn	Ser		Lvs	Asp	Asn	Ara	Glv	Asn
57	ser	тут	AIG	260	Gry	мэр	1111	111	265	1,5	2,5	op		270	U-1	
58	T	T 011	C15		т1.	CTTC	Thr	C137		G1v	Δrσ	G1 v	Glu		Lvs	Cvs
59	Leu	ьeu		Cys	TTE	Cys	1111	280	ASII	СТУ	пту	GLY	285		L 15	O _I B
60	a 1		275	mla sa	000	1701	Gln		mh r	Cor	Cor	C1v		G1 v	Dro	Dhe
61	GIU	-	HIS	THE	Ser	Val	295	1111	1 111	Ser	Ser	300	Ser	GLY	110	1110
62		290	** . 1			3 1 <u>-</u>	Val	M	Cln	Dwo	cln		uic	Dro	Cln	Dro
63		Asp	vaı	Arg	Ата		vai	TAL	GIII	PIO	315	PIU	птэ	PIO	GIII	320
64	305	_	_			310	**- 1	m\	3	a		17.0]	17-1	M	Cor	
65	Pro	Pro	Tyr	GTĀ		Cys	Val	Tnr	Asp		СТУ	vaı	Val	тут		vaı
66					325	_	_,		a 7	330	.	01	1/-4	T	335	mh m
67	Gly	Met	Gln		Leu	Lys	Thr	GIn		Asn	гĀг	GIN	мет		Cys	THE
68				340					345	-				350	a 1.	ml
69	Cys	Leu	Gly	Asn	Gly	Val	Ser		Gln	Glu	Thr	Ala		Thr	GIN	Tnr
70			355					360					365			_
71	Tyr	Gly	Gly	Asn	Ser	Asn	Gly	Glu	Pro	Cys	Val		Pro	Phe	Thr	\mathtt{Tyr}
72		370					375					380				
73	Asn	Asp	Arg	Thr	Asp	Ser	Thr	Thr	Ser	Asn		Glu	Gln	Asp	Gln	Lys
74	385					390					395					400
75	Tyr	Ser	Phe	Cys	Thr	Asp	His	Thr	Val	Leu	Val	Gln	Thr	Arg	Gly	Gly
76					405					410					415	
77	Asn	Ser	Asn	Gly	Ala	Leu	Cys	His	Phe	Pro	Phe	Leu	Tyr	Asn	Asn	His
78				420					425					430		
79	Asn	Tyr	Thr	Asp	Cys	Thr	Ser	Glu	Gly	Arg	Arg	Asp	Asn	Met	Lys	\mathtt{Trp}
80		•	435	-	-			440					445			
81	Cvs	Glv	Thr	Thr	Gln	Asn	Tyr	Asp	Ala	Asp	Gln	Lys	Phe	Gly	Phe	Cys
82		450					455	-				460				
83	Pro	Met	Ala	Ala	His	Glu	Glu	Ile	Cys	Thr	Thr	Asn	Glu	Gly	Val	Met
84	465					470			_		475					480
85	Tvr	Arg	Ile	Gly	Asp	Gln	Trp	Asp	Lys	Gln	His	Asp	Met	Gly	His	Met
86	-1-	3		•	485		-	-	-	490					495	
87	Met	Ara	Cvs	Thr	Cvs	Val	Gly	Asn	Gly	Arq	Gly	Glu	Trp	Thr	Cys	Ile
88		5	-1-	500	- 4 -		-		505	_	-		_	510	_	
89	Δla	Tvr	Ser		Leu	Ara	Asp	Gln	Cvs	Ile	Val	Asp	Asp	Ile	Thr	Tyr
90	1114	-1-	515			5		520	- 1			-	525			_
91	Δen	Va 1		Asp	Thr	Phe	His		Ara	His	Glu	Glu	Glv	His	Met	Leu
92	USII	530	21511	шь		1 110	535	2,2	5			540	1			
93	λen		Thr	Cvs	Dhe	Glv	Gln	Glv	Ara	Glv	Ara		Lvs	Cvs	Asp	Pro
94	545	_	1111	CID	1 110	550	0111			0-1	555		-1-	-1-		560
			Cln	Cvc	Cln		Sar	Glu	Thr	Glv		Dhe	Tur	Gln	Tle	Gly
95	Val	ASP	GIH	Суз	565	мэр	PCI	GIU	1111	570	1111	1110	- 1 -	0111	575	011
96	1	C	m~~	C1.,		Mirr	Val	uic	Clv		λνα	ጥነሪጉ	Gln	Cve		Cvs
97	ASP	ser	пр		гуѕ	тут	vai	птэ	585	vai	лту	1 7 1	GIII	590	- 7 -	CyD
98		a1	3	580	τĺα	<i>α</i> 1	<i>α</i> 1	m~~		Crro	Cln	Dro	LOU		Thr	ጥህኮ
99	туг	GTĀ	_		тте	стХ	GIU			Cys	GTII	FIO	60:		1111	Tyr
100	_		59		. 07	. n	. 17-	60		1 101-	A T1	o արև			r Dr	0 602
101	Pr			r se	r G1	y Pr			u Val.	r Pu	e 11	e Th.		u III.	L PI	o Ser
102		61	U				61	J				02	U			

RAW SEQUENCE LISTING DATE: 04/12/2002

PATENT APPLICATION: US/09/581,651A TIME: 14:21:03

Input Set : A:\350013-72.txt

```
Gln Pro Asn Ser His Pro Ile Gln Trp Asn Ala Pro Gln Pro Ser His
103
                                             635
104
                         630
     Ile Ser Lys Tyr Ile Leu Arg Trp Arg Pro Val Ser Ile Pro Pro Arg
105
                                         650
106
     Asn Leu Gly Tyr
107
108
                 660
110 <210> SEQ ID NO: 2
111 <211> LENGTH: 2147
112 <212> TYPE: DNA
113 <213> ORGANISM: Human
115 <400> SEQUENCE: 2
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                                                                              60
                                                                             120
     ttaggggtcc ggggcccggg ctgctgctgc tggccgtcca gtgcctgggg acagcggtgc
117
     cctccacggg agcctcgaag agcaagaggc aggctcagca aatggttcag ccccagtccc
                                                                             180
118
                                                                             240
     cqqtqqctqt caqtcaaaqc aaqcccqqtt gttatgacaa tggaaaacac tatcagataa
119
     atcaacaqtq qqaqcqqacc tacctagqca atgcgttggt ttgtacttgt tatggaggaa
                                                                             300
120
                                                                             360
     gccgaggttt taactgcgag agtaaacctg aagctgaaga gacttgcttt gacaagtaca
121
     ctgggaacac ttaccgagtg ggtgacactt atgagcgtcc taaagactcc atgatctggg
                                                                             420
122
     actgtacctg catcggggct gggcgaggga gaataagctg taccatcgca aaccgctgcc
                                                                             480
123
                                                                             540
     atgaaggggg tcagtcctac aagattggtg acacctggag gagaccacat gagactggtg
124
     gttacatgtt agagtgtgtg tgtcttggta atggaaaagg agaatggacc tgcaagccca
                                                                             600
125
     tagctgagaa gtgttttgat catgctgctg ggacttccta tgtggtcgga gaaacgtggg
                                                                             660
126
     agaagcccta ccaaggctgg atgatggtag attgtacttg cctgggagaa ggcagcggac
                                                                             720
127
                                                                             780
     gcatcacttg cacttctaga aatagatgca acgatcagga cacaaggaca tcctatagaa
128
     ttggagacac ctggagcaag aaggataatc gaggaaacct gctccagtgc atctgcacag
                                                                             840
129
     gcaacggccg aggagagtgg aagtgtgaga ggcacacctc tgtgcagacc acatcgagcg
                                                                             900
130
                                                                             960
     gatetggccc etteacegat gttegtgeag etgtttacea acegeageet eaceeceage
131
                                                                            1020
     ctcctcccta tgqccactqt qtcacagaca gtggtgtggt ctactctgtg gggatgcagt
                                                                            1080
133
     ggctgaagac acaaggaaat aagcaaatgc tttgcacgtg cctgggcaac ggagtcagct
     gccaagagac agctgtaacc cagacttacg gtggcaactc aaatggagag ccatgtgtct
                                                                            1140
134
                                                                            1200
135
     taccattcac ctacaacgac aggacggaca gcacaacttc gaattatgag caggaccaga
                                                                            1260
136
     aatactcttt ctgcacagac cacactgttt tggttcagac tcgaggagga aattccaatg
     gtgccttgtg ccacttcccc ttcctataca acaaccacaa ttacactgat tgcacttctg
                                                                            1320
137
                                                                            1380
     agggcagaaq agacaacatg aagtggtgtg ggaccacaca gaactatgat gccgaccaga
138
                                                                            1440
139
     agtttqqqtt ctqccccatq qctqcccacq aqqaaatctq cacaaccaat gaaggggtca
140
     tgtaccqcat tggagatcag tgggataagc agcatgacat gggtcacatg atgaggtgca
                                                                            1500
     cgtgtgttgg gaatggtcgt ggggaatgga catgcattgc ctactcgcag cttcgagatc
                                                                            1560
141
                                                                            1620
     agtgcattgt tgatgacatc acttacaatg tgaacgacac attccacaag cgtcatgaag
142
     aggggcacat gctgaactgt acatgcttcg gtcagggtcg gggcaggtgg aagtgtgatc
                                                                            1680
143
144
     ccgtcgacca atgccaggat tcagagactg ggacgtttta tcaaattgga gattcatggg
                                                                            1740
     agaagtatgt gcatggtgtc agataccagt gctactgcta tggccgtggc attggggagt
                                                                            1800
145
     ggcattgcca acctttacag acctatccaa gctcaagtgg tcctgtcgaa gtatttatca
                                                                            1860
     ctgagactec gagteageec aacteecace ceatecagtg gaatgeacea cagecatete
147
                                                                            1920
     acatttccaa gtacattctc aggtggagac ctgtgagtat cccacccaga aaccttggat
                                                                            1980
148
     actgagtctc ctaatcttat caattctgat ggtttctttt tttcccagct tttgagccaa
                                                                            2040
149
     caactctgat taactattcc tatagcattt actatatttg tttagtgaac aaacaatatg
                                                                            2100
150
    tggtcaatta aattgacttg tagactgaaa aaaaaaaaa aaaaaaa
                                                                            2147
153 <210> SEQ ID NO: 3
154 <211> LENGTH: 20
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RAW SEQUENCE LISTING DATE: 04/12/2002 PATENT APPLICATION: US/09/581,651A TIME: 14:21:03

Input Set : A:\350013-72.txt

```
155 <212> TYPE: PRT
156 <213> ORGANISM: Human
158 <400> SEQUENCE: 3
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                                          10
160
161 Asn Leu Gly Tyr
162
                 20
164 <210> SEQ ID NO: 4
165 <211> LENGTH: 21
166 <212> TYPE: PRT
167 <213> ORGANISM: Human
169 <400> SEQUENCE: 4
170 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Ala Leu Val Cys Thr Cys
                                          10
171
     Tyr Gly Gly Ser Arg
173
                 20
175 <210> SEQ ID NO: 5
176 <211> LENGTH: 23
177 <212> TYPE: PRT
178 <213> ORGANISM: Human
180 <400> SEQUENCE: 5
181 Pro Cys Val Leu Pro Phe Thr Tyr Asn Asp Arg Thr Asp Ser Thr Thr
     Ser Asn Tyr Glu Gln Asp Gln
183
184
                 20
186 <210> SEQ ID NO:
187 <211> LENGTH: 20
188 <212> TYPE: PRT
189 <213> ORGANISM: Human
191 <400> SEQUENCE: 6
     Thr Asp His Thr Val Leu Val Gln Thr Arg Gly Gly Asn Ser Asn Gly
                      5
                                          10
193
     1
194
    Ala Leu Cys His
195
197 <210> SEQ ID NO: 7
198 <211> LENGTH: 21
199 <212> TYPE: PRT
200 <213> ORGANISM: Human
202 <400> SEQUENCE: 7
    Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Ile Ala Tyr Ser Gln Leu
203
204
     1
205
    Arg Asp Gln Cys Ile
206
                 20
208 <210> SEQ ID NO: 8
209 <211> LENGTH: 21
210 <212> TYPE: PRT
211 <213> ORGANISM: Human
213 <400> SEQUENCE: 8
214 Gln Gln Trp Glu Arg Thr Tyr Leu Gly Asn Val Leu Val Cys Thr Cys
```

RAW SEQUENCE LISTING DATE: 04/12/2002 PATENT APPLICATION: US/09/581,651A TIME: 14:21:03

Input Set : A:\350013-72.txt

```
10
                                                              15
216 Tyr Gly Gly Ser Arg
                 20
217
219 <210> SEQ ID NO: 9
220 <211> LENGTH: 39
221 <212> TYPE: PRT
222 <213> ORGANISM: Human
224 <400> SEQUENCE: 9
225 Glu Pro Cys Val Leu Pro Phe Thr Tyr Asn Gly Arg Thr Phe Tyr Ser
226
                                          10
     Cys Thr Thr Glu Gly Arg Gln Asp Gly His Leu Trp Cys Ser Thr Thr
227
228
                 20
                                      25
229
    Ser Asn Tyr Glu Gln Asp Gln
230
             35
232 <210> SEQ ID NO: 10
233 <211> LENGTH: 21
234 <212> TYPE: PRT
235 <213> ORGANISM: Human
237 <400> SEQUENCE: 10
238 Cys Thr Asp His Thr Val Leu Val Gln Thr Gln Gly Gly Asn Ser Asn
                      5
                                          10
239
    1
240 Gly Ala Leu Cys His
241
243 <210> SEQ ID NO: 11
244 <211> LENGTH: 21
245 <212> TYPE: PRT
246 <213> ORGANISM: Human
248 <400> SEQUENCE: 11
249 Val Gly Asn Gly Arg Gly Glu Trp Thr Cys Tyr Ala Tyr Ser Gln Leu
                     5
251 Arg Asp Gln Cys Ile
252
                 20
254 <210> SEQ ID NO: 12
255 <211> LENGTH: 20
256 <212> TYPE: PRT
257 <213> ORGANISM: Human
259 <400> SEQUENCE: 12
260 Ile Ser Lys Tyr Ile Leu Arg Trp Arg Pro Lys Asn Ser Val Gly Arg
                                          10
261
262
    Trp Lys Glu Ala
263
265 <210> SEQ ID NO: 13
266 <211> LENGTH: 11
267 <212> TYPE: PRT
268 <213> ORGANISM: Human
270 <400> SEQUENCE: 13
271 Thr Ala Ser Gly Val Ala Glu Thr Thr Asn Cys
272
    1
                      5
274 <210> SEQ ID NO: 14
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VERIFICATION SUMMARY

DATE: 04/12/2002

PATENT APPLICATION: US/09/581,651A TIME: 14:21:04

Input Set : A:\350013-72.txt